



**284 Sheffield Street
Mountainside, NJ 07092**

SDG NARRATIVE

USEPA

SDG # MYDBD4

CASE # 51772

CONTRACT # 68HERH20D0011

SOW# SFAM01.1

LAB NAME: Alliance Technical Group, LLC

LAB CODE: ACE

LAB ORDER ID # P4332

MODIFIED ANALYSIS #3225.1, 3226.1

A. Number of Samples and Date of Receipt

20 Soil samples were delivered to the laboratory intact on 10/07/2024.

B. Parameters

Test requested for Metals CLP FULL = Aluminum, Antimony, Arsenic, Barium, Beryllium, Cadmium, Calcium, Chromium, Cobalt, Copper, Iron, Lead, Magnesium, Manganese, Nickel, Potassium, Selenium, Silver, Sodium, Thallium, Vanadium, Zinc.

Test requested for Metals CLP MS FULL = Antimony, Arsenic, Barium, Beryllium, Cadmium, Chromium, Cobalt, Copper, Lead, Nickel, Selenium, Silver, Thallium, Vanadium, Zinc.

C. Cooler Temp

Indicator Bottle: Presence/Absence

Cooler: 21.8°C

D. Detail Documentation (related to Sample Handling Shipping, Analytical Problem, Temp of Cooler etc):

Issue 1 : A "P" or "M" prefix was listed at the beginning of a CLP sample ID.

E. Corrective Action taken for above:

Resolution 1 : To maintain COC integrity, ASB requests no changes to the Sample IDs. The laboratory will note the issue in the SDG Narrative and proceed with the analysis of the samples.

F. Analytical Techniques:

All analyses were based on CLP Methodology by method SFAM01.1.



**284 Sheffield Street
Mountainside, NJ 07092**

Inter Element correction factors (IECs) are determined annually and correction factor are applied during ICP-AES analysis.

G. Calculation:

Calculation for ICP-AES Soil Sample:

Conversion of Results from mg/L or ppm to mg/kg (Dry Weight Basis):

$$\text{Concentration (mg/kg)} = C \times \frac{V_f}{W \times S} \times DF$$

Where,

C = Instrument value in ppm (The average of all replicate exposures)

V_f = Final digestion volume (mL)

W = Initial aliquot amount (g) (Sample amount taken in prep)

S = % Solids / 100 (Fraction of Percent Solids)

DF = Dilution Factor

Example Calculation For Sample MYDBD4 For Antimony:

If C = 0.0695554 ppm

V_f = 100 ml

W = 1.15g

S = 0.966(96.6/100)

DF = 1

$$\text{Concentration (mg/kg)} = 0.0695554 \times \frac{100}{1.15 \times 0.966} \times 1$$

$$= 12.52235 \text{ mg/kg}$$

$$= 13 \text{ mg/kg (Reported Result with Signification)}$$

Calculation for ICP-MS Soil Sample:

Conversion of Results from µg /L or ppb to mg/kg :

$$\text{Concentration (mg/kg)} = C \times \frac{V_f}{W \times S} \times DF / 1000$$

Where,

C = Instrument value in ppb (The average of all replicate integrations)

V_f = Final digestion volume (mL)

W = Initial aliquot amount (g) (Fraction of Sample amount taken in prep)



**284 Sheffield Street
Mountainside, NJ 07092**

S = % Solids / 100 (Fraction of Percent Solids)
DF = Dilution Factor

Example Calculation For Sample MYDBD4 For Antimony :

If C = 32.15 ppb

Vf = 500 ml

W = 1.15 g

S = 0.966 (96.6/100)

DF = 1

$$\text{Concentration (mg/kg)} = 32.15 \times \frac{500}{1.15 \times 0.966} \times 1 / 1000$$

$$= 14.47025 \text{ mg/kg}$$

$$= 15 \text{ mg/kg (Reported Result with Signification)}$$

H. QA/ QC

Calibrations met requirements. Interference check met requirements. Blank analyses did not indicate any presence of contamination. Laboratory Control sample was within control limits. Spike sample did meet requirements except for Antimony, Chromium, Selenium. Spike sample (MYDBE6SRE) did meet requirements except for Arsenic, Selenium, Silver, Zinc. Spike sample (MYDBE6S) did meet requirements except for Beryllium, Chromium, Selenium. Duplicate sample did meet requirements except for Antimony, Arsenic, Barium, Beryllium, Cadmium, Chromium, Cobalt, Copper, Lead, Nickel, Selenium, Silver, Vanadium, Zinc. Serial Dilution did meet requirements.

Internal standard 209Bi(1) was out Side qc limit for samples MYDBD4, MYDBD5, MYDBD7, MYDBE1, MYDBE4, MYDBE5, MYDBF1 in Original so for these samples affected parameters are reported from 2X Dilution.

Internal standard 209Bi(1) was out Side qc limit for samples MYDBD6, MYDBD8, MYDBF3 in Original & 2X Dilution so for these samples affected parameters are reported from Original.

Collision cell is being used to remove potential interferences. The analytes Na, Mg, Al, K, V, Cr, Mn, Fe, Co, Ni, Cu, Zn, As are being analyzed with collision cell and analytes Be, B, Ca, Ti, Se, Sr, Zr, Mo, Ag, Cd, Sn, Sb, Ba, Tl, Pb, U are being analyzed with Non-Collision Cell. Helium gas is used for the Collision Cell analysis.



**284 Sheffield Street
Mountainside, NJ 07092**

Internal Standard Association for ICP-MS analysis.

| Target Analyte | Associated Internal Standard |
|----------------|------------------------------|
| Antimony | 159Tb |
| Arsenic | 89Y |
| Barium | 159Tb |
| Beryllium | 6Li |
| Cadmium | 159Tb |
| Chromium | 45Sc |
| Cobalt | 45Sc |
| Copper | 45Sc |
| Lead | 209Bi |
| Nickel | 45Sc |
| Selenium | 89Y |
| Silver | 159Tb |
| Thallium | 209Bi |
| Vanadium | 45Sc |
| Zinc | 45Sc |

I certify that the data package is in compliance with the terms and conditions of the contract both technically and for completeness, for other than the conditions detailed above. Release of the data contained in this hard copy data package has been authorized by the Laboratory Director or his designee, as verified by the following signature.

Signature_____

Name: Nimisha Pandya

Date _____

Title: Document Control Officer

| | | |
|--|-----------------------|---|
| Date: 09/11/2024 | MA: 3225.1 | Title: ICP-MS with Modified Preparation Method and Analysis of Soils with Additional Laboratory QC |
| Method Source: SFAM01.1 | Method: ICP-MS | |
| Matrix: Soil/Sediment | | |
| Summary of Modification | | |
| <p>The purpose of this modified analysis is to prepare samples by EPA Draft Method 3050C (see below) with additional modified LCS and Matrix Spikes and analyze for the scheduled target analytes by ICP-MS. Unless specifically modified by this modification, all analyses, Quality Control (QC), and reporting requirements specified in the SOW listed in your current EPA agreement remain unchanged and in full force and effect.</p> | | |
| I. Analyte Modifications | | Not applicable <input checked="" type="checkbox"/> |
| II. Calibration and QC Requirements | | Not applicable <input type="checkbox"/> |
| <p>The Laboratory shall:</p> <ul style="list-style-type: none"> • Use the Method Detection Limits (MDLs) determined for routine soil analyses (i.e., Method 200.8) to report the results for these analyses. The Laboratory is NOT required to perform an MDL study for Draft Method 3050C. • Prepare and analyze an additional Laboratory Control Sample (LCS) spiked at the CRQL. Percent Recovery limits do NOT apply to this LCS and no corrective actions are required. • Prepare a Matrix Spike spiked at three times the levels specified in the SOW. • Prepare and analyze an additional Matrix Spike sample spiked at five times the levels specified for this Modified Analysis (i.e., 15x the levels specified in the SOW). • Post-Digestion Spike requirements apply to the 5x Matrix Spike only. • Post-Digestion Spike corrective actions apply to Sb. | | |
| III. Preparation and Method Modifications | | Not applicable <input type="checkbox"/> |
| <p>The Laboratory shall:</p> <ul style="list-style-type: none"> • Prepare and analyze the sample by EPA Draft Method 3050C as follows: <ul style="list-style-type: none"> ○ Mix sample thoroughly and transfer 1.00 – 1.50 g to a digestion vessel. ○ Add 10 mL 1:1 HNO₃ and 5 mL 1:1 HCl, heat the sample at 95°C (±3°C) and reflux 10-15 minutes. ○ Add 5 mL concentrated HNO₃ and reflux for 30 minutes at 95°C (±3°C), repeat until digestion complete. ○ Concentrate sample to 5 mL or reflux without boiling for 2 hours at 95°C (±3°C). ○ Cool sample, add 2mL water and 3 mL 30% H₂O₂. Heat at 95°C (±3°C) and add additional 1 mL aliquots of 30% H₂O₂ until effervescence is minimal. ○ Reduce volume to 5 mL or reflux without boiling for 2 hours at 95°C (±3°C). ○ Dilute to 100 mL with water, centrifuge or filter as necessary prior to analysis. • The same sample extracts can be used for ICP-AES analysis. Separate Matrix Spikes and LCS will need to be prepared for both ICP-AES and ICP-MS analyses. • Analyze the samples starting at an initial 5x dilution. Subsequently, dilute samples as necessary to bring the analyte concentrations within the calibration range of the instrument per the SOW. • Method Blanks, both LCSs, and all instrument QC are to be analyzed undiluted. | | |

IV. Special Reporting Requirements**Not applicable**☐

The Laboratory shall:

- Ensure the SDG Narrative is updated as stated in the SOW, including any technical and administrative problems encountered and the resolution or corrective actions taken. These problems may include interference problems encountered during analysis, dilutions, re-analyses and/or re-preparations performed, and problems with the analysis of samples. Also include a discussion of any SOW Modified Analyses, including a copy of the approved modification form with the SDG Narrative.
- Initial analysis data are reported with a dilution factor of 1.0 and a final volume of 500 mL, per the SOW.
- Report the additional LCS as "LCSD" in the raw data and in the EDD with QCType "Laboratory_Control_Sample_Duplicate".
- Report the additional Matrix Spike with an "SRE" suffix in the raw data and EDD.
- Report any Post-Digestion Spike of the additional 5x Matrix Spike with an "ARE" suffix.

| | | |
|---|------------------------|--|
| Date: 09/11/2024 | MA: 3226.1 | Title: ICP-AES with Modified Preparation Method and Analysis of Soils with Additional Laboratory QC |
| Method Source: SFAM01.1 | Method: ICP-AES | |
| Matrix: Soil/Sediment | | |
| Summary of Modification | | |
| <p>The purpose of this modified analysis is to prepare samples by EPA Draft Method 3050C (see below) with additional modified LCS and Matrix Spikes and analyze for the scheduled target analytes by ICP-AES. Unless specifically modified by this modification, all analyses, Quality Control (QC), and reporting requirements specified in the SOW listed in your current EPA agreement remain unchanged and in full force and effect.</p> | | |
| I. Analyte Modifications | | Not applicable <input checked="" type="checkbox"/> |
| II. Calibration and QC Requirements | | Not applicable <input type="checkbox"/> |
| <p>The Laboratory shall:</p> <ul style="list-style-type: none"> • Use the Method Detection Limits determined for routine soil analyses (i.e., Method 3050B) to report the results for these analyses. The Laboratory is NOT required to perform an MDL study for Draft Method 3050C. • Prepare and analyze an additional Laboratory Control Sample (LCS) spiked at the CRQL. Percent Recovery limits do NOT apply to this LCS and no corrective actions are required. • Prepare a Matrix Spike spiked at two times the levels specified in the SOW. • Post-Digestion Spike requirements apply to the 2x Matrix Spike. • Post-Digestion Spike corrective actions apply to Sb. | | |
| III. Preparation and Method Modifications | | Not applicable <input type="checkbox"/> |
| <p>The Laboratory shall:</p> <ul style="list-style-type: none"> • Prepare and analyze the sample by EPA Draft Method 3050C as follows: <ul style="list-style-type: none"> ○ Mix sample thoroughly and transfer 1.00 – 1.50 g to a digestion vessel. ○ Add 10 mL 1:1 HNO₃ and 5 mL 1:1 HCl, heat the sample at 95°C (±3°C) and reflux 10 -15 minutes. ○ Add 5 mL concentrated HNO₃ and reflux for 30 minutes at 95°C (±3°C), repeat until digestion complete. ○ Concentrate sample to 5 mL or reflux without boiling for 2 hours at 95°C (±3°C). ○ Cool sample, add 2mL water and 3 mL 30% H₂O₂. Heat at 95°C (±3°C) and add additional 1 mL aliquots of 30% H₂O₂ until effervescence is minimal. ○ Reduce volume to 5 mL or reflux without boiling for 2 hours at 95°C (±3°C). ○ Dilute to 100 mL with water, centrifuge or filter as necessary prior to analysis. • The same sample extracts can also be used for ICP-MS analysis. Separate Matrix Spikes and LCS will need to be prepared for both ICP-AES and ICP-MS analyses. • Analyze the samples starting at an initial 2x dilution. Subsequently, dilute samples as necessary to bring the analyte concentrations within the calibration range of the instrument per the SOW. • Verify that the dilution was adequate to reduce interferences to within the method calibration range. This can optionally be verified by visual verification of the spectrogram or by analysis of a serial dilution. There are other acceptable means to provide assurance, e.g. some software may automatically provide guidance to the analyst. • Method Blanks, both LCS, and all instrument QC are to be analyzed undiluted. | | |

| | |
|---|--|
| | |
| IV. Special Reporting Requirements | Not applicable <input type="checkbox"/> |
| <p>The Laboratory shall:</p> <ul style="list-style-type: none">• Ensure the SDG Narrative is updated as stated in the SOW, including any technical and administrative problems encountered and the resolution or corrective actions taken. These problems may include interference problems encountered during analysis, dilutions, re-analyses and/or re-preparations performed, and problems with the analysis of samples. Also include a discussion of any SOW Modified Analyses, including a copy of the approved modification form with the SDG Narrative.• Initial analysis data are reported with a dilution factor of 2.0 and a final volume of 100 mL, per the SOW.• Report the additional LCS as "LCSD" in the raw data and in the EDD with QCType "Laboratory_Control_Sample_Duplicate".• Ensure that up-to-date Interelement Correction Factors (IECs) are provided with the data package. | |

| Element, Wavelength and Order | Use? | # IECs | IEC | k1 | k2 | Calc-in-fit? |
|-------------------------------------|-------------------------------------|--------|-----|-----------|----------|--------------|
| As 189.042 {479} | <input checked="" type="checkbox"/> | 1 | Fe | -0.000064 | 0.000000 | No |
| Ti 190.856 {477} | <input checked="" type="checkbox"/> | 5 | Mo | -0.002450 | 0.000000 | No |
| | | | Co | 0.002248 | 0.000000 | No |
| | | | Ti | -0.000500 | 0.000000 | No |
| | | | Mn | 0.000370 | 0.000000 | No |
| | | | V | -0.012340 | 0.000000 | No |
| Pb 220.353 {453} | <input checked="" type="checkbox"/> | 6 | Mo | -0.001480 | 0.000000 | No |
| | | | Al | -0.000075 | 0.000000 | No |
| | | | Cu | 0.001400 | 0.000000 | No |
| | | | Fe | 0.000030 | 0.000000 | No |
| | | | Mn | 0.000340 | 0.000000 | No |
| | | | Ni | 0.000630 | 0.000000 | No |
| Se 196.090 {472} | <input checked="" type="checkbox"/> | 3 | Fe | -0.000308 | 0.000000 | No |
| | | | Mn | 0.000470 | 0.000000 | No |
| | | | Co | -0.000630 | 0.000000 | No |
| Sb 206.833 {463} | <input checked="" type="checkbox"/> | 4 | Cr | 0.010700 | 0.000000 | No |
| | | | V | -0.001168 | 0.000000 | No |
| | | | Mo | -0.002850 | 0.000000 | No |
| | | | Ni | -0.000440 | 0.000000 | No |
| Al 396.152 { 85} | <input checked="" type="checkbox"/> | 1 | Mo | 0.037230 | 0.000000 | No |
| Ba 493.409 { 68} | <input type="checkbox"/> | None | | | | |
| Be 234.861 {144} | <input checked="" type="checkbox"/> | 3 | Mo | -0.000320 | 0.000000 | No |
| | | | Fe | 0.000010 | 0.000000 | No |
| | | | Mn | -0.000047 | 0.000000 | No |
| Cd 214.438 {457} | <input checked="" type="checkbox"/> | 1 | Fe | 0.000040 | 0.000000 | No |
| Ca 373.690 { 90} | <input type="checkbox"/> | None | | | | |
| Cr 267.716 {126} | <input checked="" type="checkbox"/> | 1 | Mn | 0.000160 | 0.000000 | No |
| Co 228.616 {448} | <input checked="" type="checkbox"/> | 2 | Ti | 0.001840 | 0.000000 | No |
| | | | Mo | -0.001230 | 0.000000 | No |
| Cu 324.754 {104} | <input checked="" type="checkbox"/> | 4 | Co | -0.000796 | 0.000000 | No |
| | | | Fe | -0.000100 | 0.000000 | No |
| | | | Mn | 0.000345 | 0.000000 | No |
| | | | Ni | 0.000895 | 0.000000 | No |
| Fe 259.837 {130} | <input type="checkbox"/> | None | | | | |
| Mn 257.610 {131} | <input checked="" type="checkbox"/> | 1 | Ni | 0.000897 | 0.000000 | No |
| Mg 279.079 {121} | <input type="checkbox"/> | None | | | | |
| Ni 231.604 {446} | <input type="checkbox"/> | None | | | | |
| Ag 328.068 {103} | <input checked="" type="checkbox"/> | 3 | Fe | -0.000100 | 0.000000 | No |
| | | | Mn | 0.000146 | 0.000000 | No |
| | | | V | -0.000889 | 0.000000 | No |
| Na 818.326 { 41} | <input type="checkbox"/> | None | | | | |
| V 292.402 {115} | <input checked="" type="checkbox"/> | 2 | Mo | -0.008480 | 0.000000 | No |
| | | | Cr | -0.002220 | 0.000000 | No |
| Zn 206.200 {464} | <input type="checkbox"/> | None | | | | |
| Zn 213.856 {158} | <input checked="" type="checkbox"/> | 1 | Ni | 0.007280 | 0.000000 | No |
| K 769.896 { 44} | <input type="checkbox"/> | None | | | | |
| P 177.495 {490} | <input checked="" type="checkbox"/> | 2 | Ni | 0.001640 | 0.000000 | No |
| | | | Cu | -0.012530 | 0.000000 | No |
| B 249.678 {135} | <input checked="" type="checkbox"/> | 3 | Co | 0.002880 | 0.000000 | No |
| | | | V | -0.002000 | 0.000000 | No |
| | | | Fe | -0.001360 | 0.000000 | No |
| Mo 202.030 {467} | <input type="checkbox"/> | None | | | | |
| S 182.034 {485} | <input checked="" type="checkbox"/> | 2 | Mo | -0.008000 | 0.000000 | No |
| | | | Mn | 0.002700 | 0.000000 | No |

| Element, Wavelength and Order | Use? | # IECs | IEC | k1 | k2 | Calc-In-fit? |
|-------------------------------------|-------------------------------------|--------|-----|-----------|----------|--------------|
| Si 251.611 {134} | <input checked="" type="checkbox"/> | 2 | Mo | 0.010520 | 0.000000 | No |
| | | | Ti | 0.005650 | 0.000000 | No |
| Sn 189.989 {478} | <input type="checkbox"/> | None | | | | |
| Ti 336.121 {100} | <input checked="" type="checkbox"/> | 1 | Ni | -0.001000 | 0.000000 | No |
| Li 670.784 { 50} | <input type="checkbox"/> | None | | | | |
| Y 224.306 {450}* | <input type="checkbox"/> | None | | | | |
| Y 360.073 { 94}* | <input type="checkbox"/> | None | | | | |
| Y 371.030 { 91}* | <input type="checkbox"/> | None | | | | |
| Y 224.306 {150}* | <input type="checkbox"/> | None | | | | |
| In 230.606 {446}* | <input type="checkbox"/> | None | | | | |
| Sr 407.771 { 83} | <input type="checkbox"/> | None | | | | |